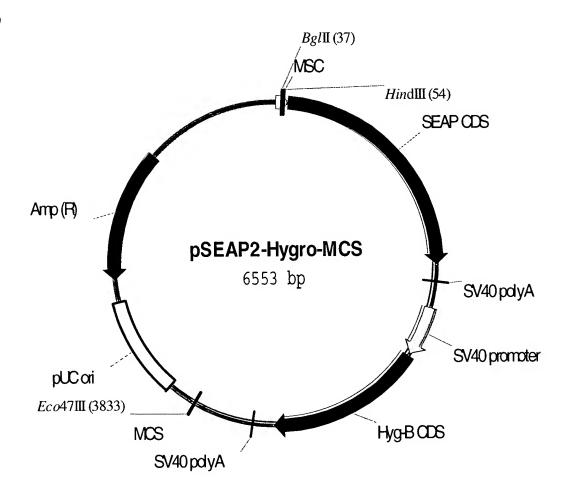
24

1/10

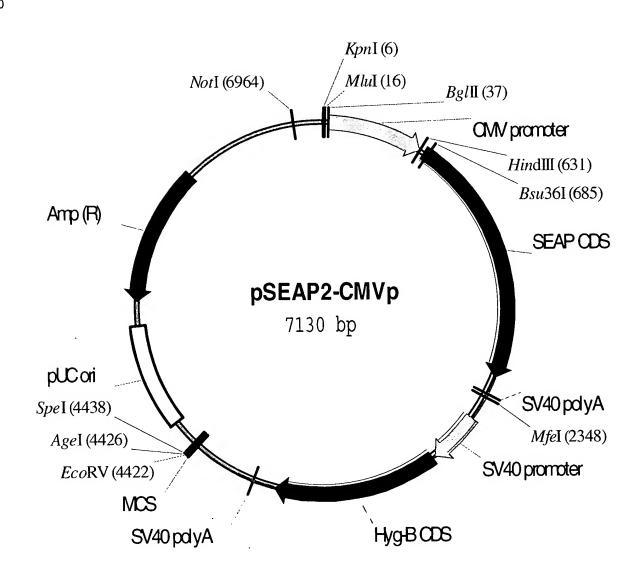
Figure 1



²⁵ 2/10

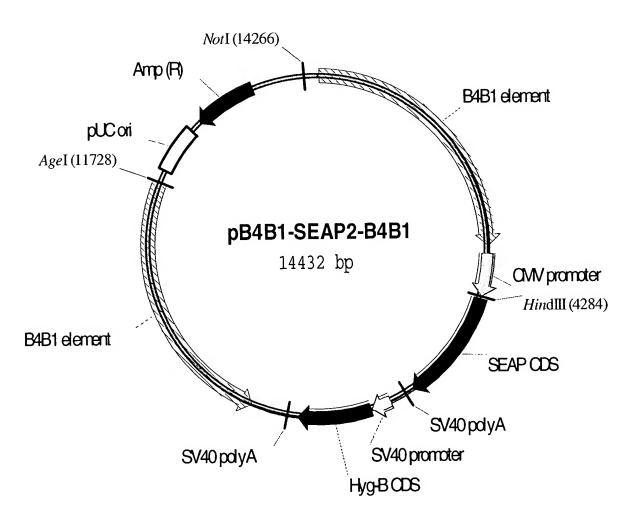
Figure 2

5



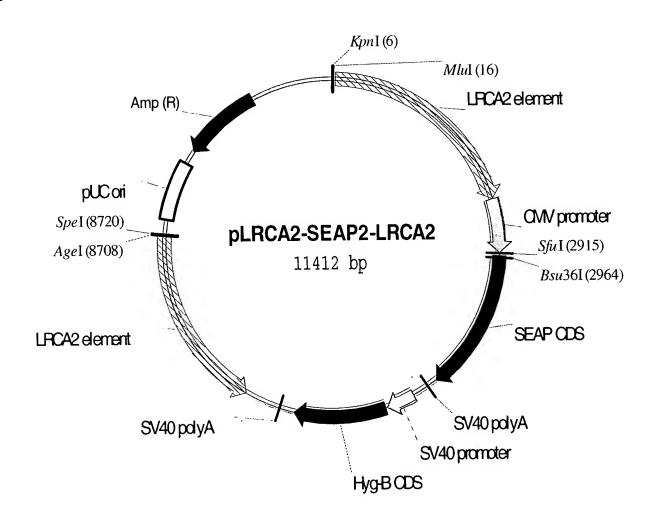
²⁶ 3/10

5 Figure 3



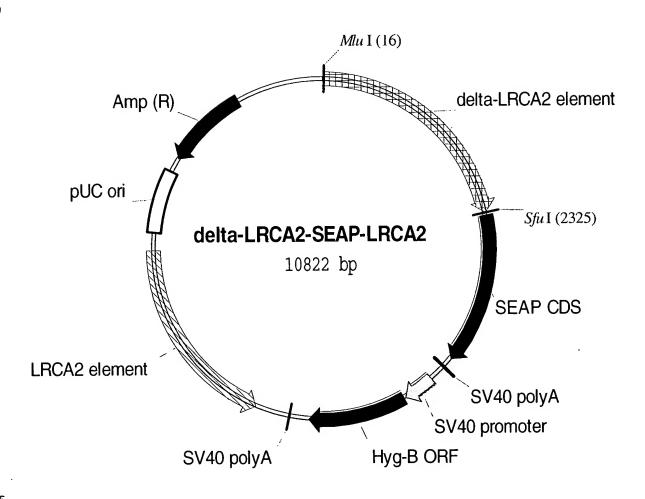
²⁷ 4/10

Figure 4



²⁸ 5/10

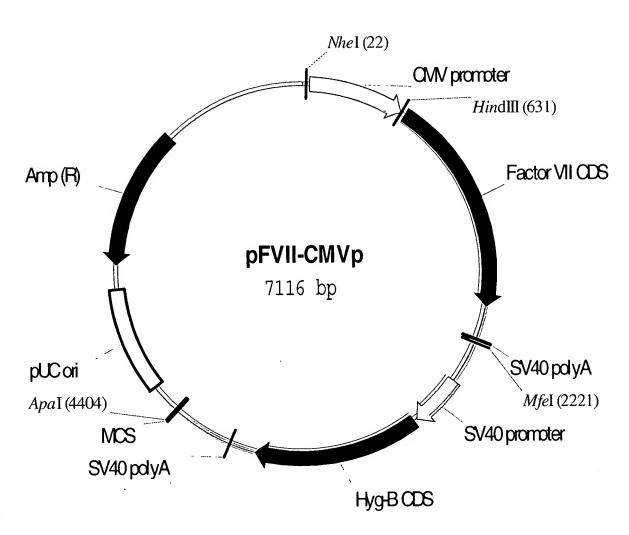
5 Figure 5



²⁹ 6/10

Figure 6

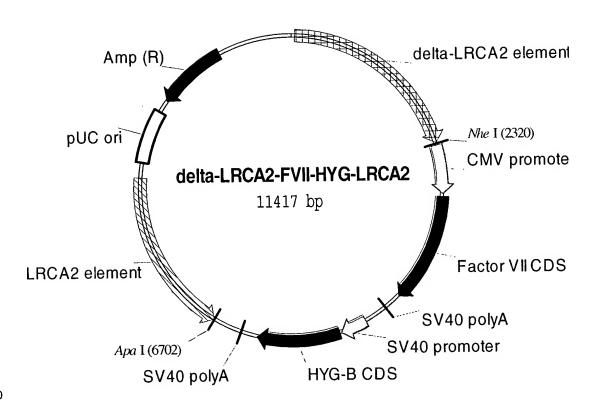
5



³⁰ 7/10

Figure 7

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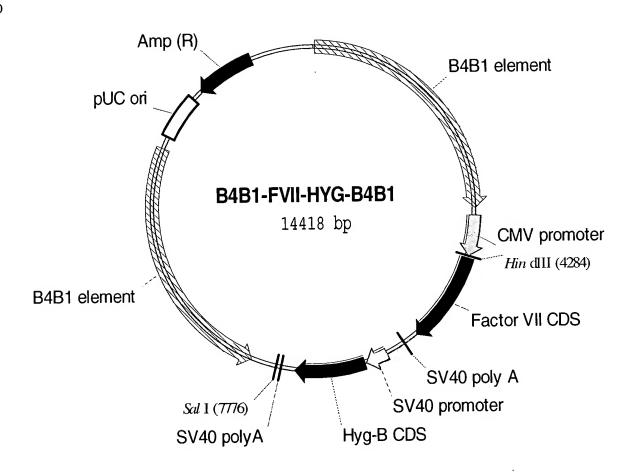


³¹ 8/10

Figure 8

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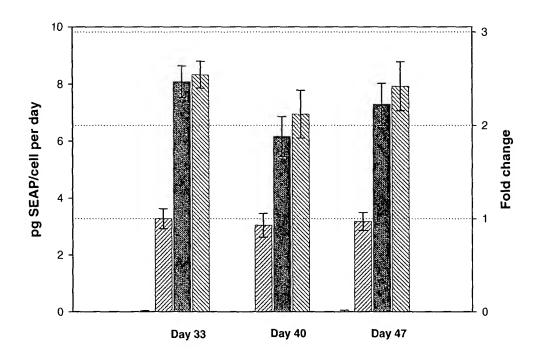


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Figure 9

5



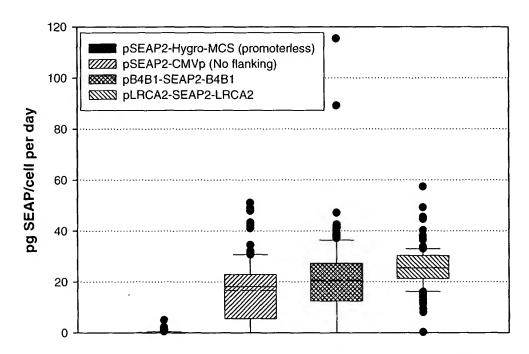
pSEAP2-Hygro-MCS (promoterless)
pSEAP2-CMVp (No flanking)
pB4B1-SEAP2-B4B1 (B4B1-flanking)
pLRCA2-SEAP2-LRCA2 (LRCA2-flanking)

33

10/10

5

Figure 10



Box plot showing statistical values calculated from single cell clone SEAP expressors. Upper and lower box boundaries indicates the 25th and 75th percentiles, respectively. Lines within boxes mark the median and dashed lines shown the mean. Whiskers above and below boxes indicate the 10th and 90th percentiles. Outlying data points are indicated by filled ovals.

Statistics were calculated from 140, 132, 144, and 128 data points, respectively.

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15